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| APPLICATION NO. | FILING DATE | FIRST NAMED INVENTOR | ATTORNEY DOCKET NO. | CONFIRMATION NO |
|-----------------|-------------------------------------|----------------------|---------------------|-----------------|
| 10/734,782 | 04/02/2004 | Steven Griffiths | VA/H-32318B | 3493 |
| 1095 | 7590 01/24/2006 | | EXAM | INER |
| NOVARTIS | | | CHEN, STAC | CY BROWN |
| | E INTELLECTUAL PRO H PLAZA 104/3 | OPERTY | ART UNIT | PAPER NUMBER |
| EAST HANO | VER, NJ 07936-1080 | | 1648 | |

DATE MAILED: 01/24/2006

Please find below and/or attached an Office communication concerning this application or proceeding.

| | Application No. | Applicant(s) |
|--|--|---|
| | 10/734,782 | GRIFFITHS ET AL. |
| Office Action Summary | Examiner | Art Unit |
| | Stacy B. Chen | 1648 |
| The MAILING DATE of this communication app Period for Reply | ears on the cover sheet with the c | orrespondence address |
| A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period w - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b). | ATE OF THIS COMMUNICATION 16(a). In no event, however, may a reply be tim 11 apply and will expire SIX (6) MONTHS from 12 cause the application to become ABANDONEI | l. ely filed the mailing date of this communication. 0 (35 U.S.C. § 133). |
| Status | | |
| 1) Responsive to communication(s) filed on 12 De | ecember 2003. | |
| ,— · | action is non-final. | |
| 3) Since this application is in condition for allowan | ice except for formal matters, pro | secution as to the merits is |
| closed in accordance with the practice under E | x parte Quayle, 1935 C.D. 11, 45 | 3 O.G. 213: |
| Disposition of Claims | | |
| 4) Claim(s) 38-42 is/are pending in the application | 1. | |
| 4a) Of the above claim(s) is/are withdraw | n from consideration. | |
| 5) Claim(s) is/are allowed. | | |
| 6)⊠ Claim(s) <u>38-42</u> is/are rejected. | | |
| 7) Claim(s) is/are objected to. | | |
| 8) Claim(s) are subject to restriction and/or | election requirement. | |
| Application Papers | | |
| 9) The specification is objected to by the Examine | r. | |
| 10)⊠ The drawing(s) filed on 02 April 2004 is/are: a) | oxtimes accepted or b) $oxtimes$ objected to t | by the Examiner. |
| Applicant may not request that any objection to the | drawing(s) be held in abeyance. See | e 37 CFR 1.85(a). |
| Replacement drawing sheet(s) including the correcti | | |
| 11) ☐ The oath or declaration is objected to by the Ex | aminer. Note the attached Office | Action or form PTO-152. |
| Priority under 35 U.S.C. § 119 | | |
| 12) Acknowledgment is made of a claim for foreign a) All b) Some * c) None of: 1. Certified copies of the priority documents 2. Certified copies of the priority documents 3. Copies of the certified copies of the prior application from the International Bureau * See the attached detailed Office action for a list of the priorical state. | s have been received. s have been received in Applicati ity documents have been receive i (PCT Rule 17.2(a)). | on No. <u>10/049,086</u> . ed in this National Stage |
| Attachment(s) | | (DTO 440) |
| Notice of References Cited (PTO-892) Notice of Draftsperson's Patent Drawing Review (PTO-948) | 4) Interview Summary Paper No(s)/Mail Da | |
| 2) ☐ Notice of Dialisperson's Fateth Brawning Neview (FFO-340) 3) ☐ Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08) Paper No(s)/Mail Date | | atent Application (PTO-152) |

Application/Control Number: 10/734,782 Page 2

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DETAILED ACTION

1. Applicant's preliminary amendment filed December 12, 2003 is acknowledged and entered. Claims 38-42 are pending and under examination.

Amendment to Correct Inventorship 37 C.F.R. 1.48(a)

2. In view of the papers filed January 18, 2005, it has been found that this nonprovisional application, as filed, through error and without deceptive intent, improperly set forth the inventorship, and accordingly, this application has been corrected in compliance with 37 CFR 1.48(a). The inventorship of this application has been changed by adding the name Joel Heppell as one of the co-inventors.

Specification

- 3. The specification is objected to because of the following:
 - ❖ The sequences in the drawings must be referenced by sequence identifiers. (SEQ ID NO:)
 - ❖ The first line of the first page of the specification fails to reflect the updated status of parent application USSN 10/049,086, now US Patent 6,919,083.
 - ❖ Figures 7 and 8 are not referenced in the specification.

Correction is required to overcome these objections to the specification.

Priority

4. Acknowledgment is made of Applicant's claim for foreign priority under 35 U.S.C. 119(a)-(d). Certified copies of 9918588.6 and 0005848.7 have been filed in parent

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Application No. 10/049,086, filed on February 6, 2002. It is noted, however, that Applicant has not filed a certified copy of the 0006674.6 application as required by 35 U.S.C. 119(b). The examiner notes that this document has been previously indicated as present in the parent application. However, a copy of the document cannot be located in the parent application. Applicant is requested to file a copy of the document. The Office regrets any inconvenience.

Claim Rejections - 35 USC § 112

5 The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 38-42 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention. The claims are drawn to an isolated protein comprising an amino acid sequence according to SEQ ID NO: 4 or a derivative thereof, wherein the derivative thereof has antigenic or immunogenic characteristics of the amino acid sequence according to SEQ ID NO: 4. Also claimed are kits, vaccines and pharmaceutical compositions comprising the claimed peptide.

The metes and bounds of the claimed peptide derivative cannot be determined without a clear definition of the derivative. While Applicant has provided a functional definition of the claimed peptide, the structural definition is lacking. It is unclear what portions of SEQ ID NO: 4 are retained in the derivative so that it remains immunogenic. Further, it is unclear what antigenic or immunogenic characteristics of the derived peptide must be shared with SEQ ID NO: 4. It is suggested that references to derivatives be deleted from the claims in order to overcome this rejection.

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Conclusion

6. The prior art made of record and not relied upon is considered pertinent to Applicant's disclosure. With regard to SEQ ID NO: 4, US Patent 6,471,964 B1, discloses and claims a sequence having 97.4% sequence identity to SEQ ID NO: 4 (see attached Sequence Alignment). Because the priority date granted to the patent (October 18, 1999) is later than the earliest priority date of the instant application (August 8, 1999), the patent is not available as prior art, although the derivatives instantly claimed are encompassed by the disclosure and claims of the patent.

No claim is allowed. Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Stacy B. Chen whose telephone number is 571-272-0896. The examiner can normally be reached on M-F (7:00-4:30). If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, James C. Housel can be reached on 571-272-0902. The fax phone number for the organization where this application or proceeding is assigned is 703-872-9306.

Stacy B. Chen

January 18, 2006

Stacy B. OL

- Sequence Alignment-

GenCore version 5.1.6 Copyright (c) 1993 - 2006 Compugen Ltd.

OM protein - protein search, using sw model

January 13, 2006, 09:27:05 ; Search time 27 Seconds (without alignments) 1886.230 Million cell updates/sec

US-10-734-782-4 3097 1 MADKGATYSFDVRDNTLVVR......DMTPRIBFDBDDBEBEBDIDI Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Scoring table:

572060 seqs, 82675679 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Total number of hits satisfying chosen parameters:

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Issued Patents AA:*

1: /cgn2 6/ptodate/1/laa/5 COMB.pep:*

2: /cgn2 6/ptodate/1/laa/f COMB.pep:*

3: /cgn2 6/ptodate/1/laa/H COMB.pep:*

4: /cgn2 6/ptodate/1/laa/H COMB.pep:*

5: /cgn2 6/ptodate/1/laa/RE COMB.pep:*

6: /cgn2 6/ptodate/1/laa/RE COMB.pep:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

| | Description | 4, | 'n | 4 | 43, | 6970 | ý | ģ | , | 4 | 69 | 65 | ď | Sequence 2, Appli | 'n | 'n | 'n | Sequence 2, Appli | 7 | 67 | G | 48 | | 79 | | 27, | ۲, | Sequence 1, Appli |
|-----------|--------------------------|-----------------|------------------|------------------|------------------|---------------------|------------------|-----------------|------------------|------------------|------------------|------------------|-----------------|-------------------|-----------------|-----------------|-----------------|-------------------|-----------------|------------------|------------------|------------------|-----------------|--------------------|--------------------|-------------------|-----------------|-------------------|
| SUMMAKIES | αı | US-10-049-086-4 | US-09-690-185A-2 | US-09-206-942-45 | US-09-206-942-43 | US-09-107-532A-6970 | US-09-077-098A-6 | US-10-192-584-6 | US-09-200-650B-7 | US-08-477-451-11 | US-09-206-942-69 | US-10-193-764-65 | US-08-038-682-2 | US-08-302-832-2 | US-08-530-198-2 | US-08-469-880-2 | US-08-728-470-2 | US-08-617-697-2 | US-08-719-641-2 | US-09-206-942-67 | US-10-193-764-63 | US-09-268-347-48 | US-09-841-786-1 | US-09-949-016-7935 | US-09-710-279-2906 | US-09-180-422B-27 | US-09-079-030-1 | US-09-108-006C-1 |
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| | Query Match Length DB | 616 | 616 | 1095 | 1101 | 564 | 2042 | 2042 | 1166 | 1781 | 1095 | 1095 | 1536 | 1536 | 1536 | 1536 | 1536 | 1536 | 1536 | 1536 | 1536 | 2048 | 3241 | 504 | 1572 | 4536 | 4536 | 4563 |
| de | Query | 100.0 | 97.4 | 4.1 | 4.1 | 3.8 | 3.8 | 3.8 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.7 | 3.6 | 3.6 | 3.6 | 3.6 | 3.6 |
| | Score | 3097 | 3015 | 128 | 128 | 118 | 116.5 | 116.5 | 116 | 116 | 115 | 115 | 115 | 115 | 115 | 115 | 115 | 115 | 115 | 115 | 115 | 114 | 114 | 113 | 112.5 | 112.5 | 112.5 | 112.5 |
| | Result No. | - | 7 | e | 4 | S | 9 | 7 | æ | σ | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 | 24 | 25 | 26 | . 27 |

| 842, | Sequence 29, Appl | Sequence 29, Appl | Sequence 4939, Ap | Sequence 4, Appli | Sequence 4, Appli | Sequence 3, Appli | Sequence 3, Appli | Sequence 29927, A | Sequence 10, Appl | Sequence 4, Appli | Sequence 57, Appl |
|-------------------|-------------------|-------------------|---------------------|-------------------|-------------------|-------------------|-------------------|----------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|
| US-09-538-092-842 | US-09-345-882-29 | US-10-071-179-29 | US-09-134-000C-4939 | US-08-621-944A-4 | US-08-945-567D-4 | US-08-621-944A-3 | US-08-945-567D-3 | US-09-252-991A-29927 | US-08-968-685A-10 | US-08-038-682-4 | US-08-302-832-4 | US-08-530-198-4 | US-08-469-880-4 | US-08-728-470-4 | US-08-617-697-4 | US-08-719-641-4 | US-09-206-942-57 |
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| 112.5 | 111.5 | 111.5 | 111 | 111 | 111 | 111 | 111 | 110.5 | 110 | 109.5 | 109.5 | 109.5 | 109.5 | 109.5 | 109.5 | 109.5 | 109 |
| 28 | 53 | 30 | 31 | 32 | 33 | 34 | 35 | 36 | 37 | 38 | 39 | 40 | 41 | 42 | 43 | 44 | 45 |

| | ; Length 616; 9; Indels 0; Gaps 0; | SLLOKAPAGTEDEFWVELDQD 60 | PSKRAAANIEWAGVDDERAGG 120 | SEGEMSIWQNGQAIIRILALA 180 | ADLKAAFROSRPKKSDYRKGQ 240 | FINASYTTLARGANIPSKYLRH 300 |
|--|--|--|---|---|---|--|
| pplication US/10049086 19083 MARTON: qua Health (Europe) Limit Griffiths, Steven EXITON: Sequence CE: P24268-/GST/RMC ICATION NUMBER: US/10/049 MIO ANTE: 2002-02-06 ATION NUMBER: GB0005848.7 DATE: 2000-03-11 DATE: 2000-03-11 ATION NUMBER: GB0005674.6 DATE: AND | / Match 100.0%; Score 3097; DB 2; Local Similarity 100.0%; Pred. No. 3.7e-309 nes 616; Conservative 0; Mismatches 0; | 1 MADKGMTYSFDYRDNTLVVRRSTATKSGIKISYREDRGTSLLOKAFAGTEDEFWVELDOD | 61 VYVDKKKRRELEBEKMKDMSTRVSGAVAAAIERSVEFDNFSKEAAANIEWAGVDDEEAGG | 121 GGLVDNRRKNKGVSNNAXNLSLFIGMVFPALTTFFSAILSEGEMSIWQNGQAIIRILALA 17 SGLVDNRRKNKGVSNNAXNLSLFIGMVFPALTTFFSAILSEGEMSIWQNGQAIIRILALA | 181 DEDGKROTRTGGORVDMADVTKLNVVTANGKVKQVEVNLANDLKAAFRQSRPKRSDVRKGQ | 241 GSKATESSISNOCMALIMKSVLSADOLFAPGVKAMRTNGFNASYTTLAEGANIPSKYLRH |
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US-09-690-185A-2
; Sequence 2, Application US/09690185A
; Patent NO. 6471964
; GENERAL INFORMATION:
; APPLICANT: BIRRING,
; APPLICANT: BIRRING,
; TITLE OF INVENTION: DNA ENCODING STRUCTURAL PROTEIN-1 OF INFECTIOUS SALMON
; TITLE OF INVENTION: DNAEMIA VIRUS AND USES THEREOF
; TILE REFERENCE: BIRRING
; CURRENT APPLICATION NUMBER: US/09/690,185A
; CURRENT APPLICATION NUMBER: EP99203401.7
; PRIOR FILING DATE: 1999-10-18
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 2
; LENGTH: 616
; TYPE: PRT
; ORGANISM: Infectious salmon anemia virus
US-09-690-185A-2.
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120 120 180 240 240 300 241 GSKATESSISNQCMALIMKSVLSADQLFAPGVKMMRTNGFNASYTTLAEGANIPSKYLRH 300 301 MRNCGGVALDLMGMKRIKNSPEGAKSKIPSIIQKKVRGRCRIEEQRLLISALKISDGENK 360 9 9 1 MADKGMTYSFDVRDNTLVVRRSTATKSGIKISYREDRGTSLLQKAFAGTDDEFWLELDQD VXVDKKI RKFLEBEKMKDMSTRVSGAVAAIERSVEFDNFSKEAAANI EMAGVDDBEAGG VYVDKRIRKFLEBEKMKDMSPRCSGSVAAAIERSVEPDNFSKEAAANIEMSGEDEBEAGG 181 DEDGKRQTRTGGQRVDMADVTKLNVVTANGKVKQVEVNLNDLKAAFRQSRPKRSDYRKGQ 241 GSKATESSISNQCMALIMKSVLSADQLFAPGVKMMRTNGFNASYTTLAEGANIPSKYLRH 1 MADKGMTYSPDVRDNTLVVRRSTATKSGIKISYREDRGTSLLQKAFAGTEDEFWVELDQD SGLVDNRRKNKGVSNMAYNLSLPIGMVFPALTTFFSAILSEGEMSIWONGQAIIRILALA DEDGKRQTRTGGQRVDMADVTKLNVVTANGKVKQVEVNLNDLKAAFRQSRPKRSDYRKGQ Gaps ö 97.4%; Score 3015; DB 2; Length 616; 96.3%; Pred. No. 1e-300; tive 17; Mismatches 6; Indels Query Match Best Local Similarity 96.3% Matches 593; Conservative 61 61 121

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280 FNASYTTLAEGANIPSKYLRHMRNCGGVALDLMGMKRIKNSPEGAKSKIFSIIQKKVRGR 339

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| 셤 | 301 MRNCGGVALDLMGMKRIKNSPEGAKSKIFSIIQKKVRGRCRTEEQRLLTSALKISDGENK 360 |
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| è | 361 FORIMDILCTSFLIDPRITKCFIPPISSLAMYIOEGNSVLAMDEMKNGEDACKICKBAK 420 |
| අ | 361 FQRIMDTLCTSFLIDPPRTTKCFIPPISSLLTYIQDGNSVLAMDFMKNGEDACRICREAK 420 |
| à | 421 LKVGVNSTFTMSVARTCVAVSMVATAFCSADIIENAVPGSERYRSNIKANTTKFKKDSTY 480 |
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| ò | 481 TIQGIRLSNVRYBARPETSQSNTDRSWQVNVTDSFGGLAVFNQGAIREMLGDGTSETTSV 540 |
| qq | 481 TIQGLELSNVKYEARPETSQSNTDRSWQVNVTDSFGGLAVFNQGAIREMLGDGTSETTSV 540 |
| Ğ | 541 NVRALVKRILKSASERSARAVKTFWVGEQGKSAIVISGVGLFSIDFEGVERAERITDMTP 600 |
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| à | 601 RIEFDEDDERERDIDI 616 |
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| US-09-20 | 42-45 |
| ; Sequence ; Patent No | ice 45, Application US/U9206942 |
| - | GENERAL INFORMATION: |
| APPLI | Yang, Yan-Ping |
| | OF INVENTION: Protective Recomb |
| | OF INVENTION: Molecular Weight Proteins |
| | 43.1 |
| ; CURRENT | |
| , EARLIER | س |
| ; NUMBER | NUMBER OF SEQ ID NOS: 95 SOPTWARE: Patentin Ver. 2.1 |
| ; SEQ ID NO | 45 |
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|) ORGP | |
| ్ర | 4.1%; Score 128; DB 2 |
| Best Lo Matches | cal Similarity 19.1%; Pred. No. 0.0023; 121; Conservative 73; Mismatches 264; |
| ò | 32 SYREDRGTSLLOKAFAGTEDEFWVELDQDVYVDKKIRKFLEBERKMKD 78 |
| đ | 332 NFNASEGSELRFKSEGSTRTAFTIESDLTLNATGGNISLNQVAGIDGNLQKSLVANKN 389 |
| ò | 79 MSTRVSGAVAAAIBASVEF-DNFSKEAAANIEMAGVDDBEAGGSGLVDNRRKNKGV 133 |
| DP QC | 390 ITFEGGNITLAADKKPIEIKGNITVKEGANVTLRSANYGNDKSALSIRGNVTN-KGNLTV 448 |
| ò | 134 SNWAYNLSKFIGMVFPALTTFFSALLSEGEMSIWQNGQAITRILALADEDGKRQTRT 190 |
| QQ | 449 TGSAINTEKNLTVEGSAKFLANPNYSFNVSGLFDNQGKSNISIAKG 494 |
| | |